

Nonreversible MCMC for discontinuous targets and disconnected spaces with boundaries

Jere Koskela

Stochastic analysis of the neutron transport equation
17 September 2021



Outline

The zig-zag process

The zig-zag process on a hybrid state space

The coalescent

A geometric embedding of coalescent trees

Zig-zag for the coalescent

One slide on neutrons

The zig-zag process¹

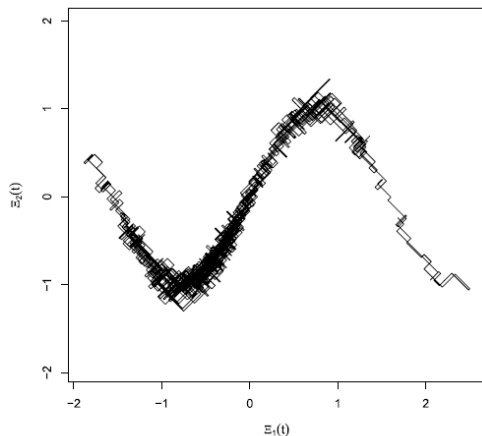
1. Set $(x, v) \in \mathbb{R}^n \times \{-1, 1\}^n$.
2. Set $t \leftarrow 0$.
3. While $t < T$ do
 - 3.1 Sample $Y \sim \text{Exp}(1)$.
 - 3.2 Set ρ such that $\sum_{i=1}^n \int_t^{t+\rho} \lambda_i(x + sv, v) ds = Y$.
 - 3.3 Set $x \leftarrow x + \rho v$.
 - 3.4 Set $t \leftarrow t + \rho$.
 - 3.5 Sample $I \sim \text{Categorical}(\lambda_1(x, v), \dots, \lambda_n(x, v))$.
 - 3.6 Set $v_I \leftarrow -v_I$.

- ▶ Between switches, x moves with constant velocity v .
- ▶ Target density π is invariant if

$$\lambda_i(x, v) = v_i \partial_i \log \pi(x) \vee 0.$$

¹J Bierkens, P Fearnhead and G Roberts. The zig-zag process and super-efficient sampling for Bayesian analysis of big data, Ann Stat 47(3):1288–1320, 2019.

The zig-zag process



(d) 2D S-shaped density

¹J Bierkens, P Fearnhead and G Roberts. The zig-zag process and super-efficient sampling for Bayesian analysis of big data, Ann Stat 47(3):1288–1320, 2019.

Advantages and disadvantages of zig-zag

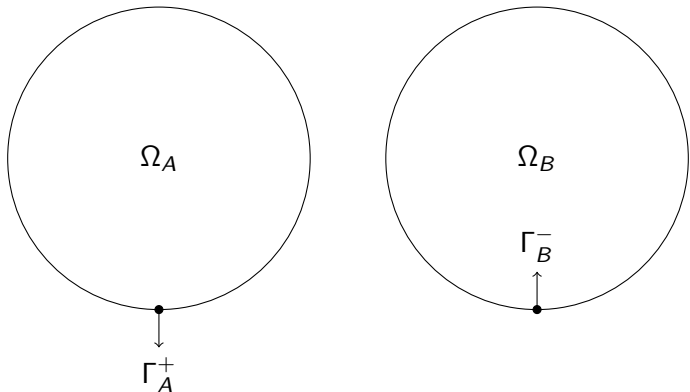
- + Avoids reversible backtracking.
- + Rejection-free.
- + Direction of travel guided by shape of target.
- Construction only makes sense if target density is differentiable. . .
- and if the state space is one connected set.

The zig-zag process on a hybrid state space

- ▶ Target $\pi(m, x)$ on $\mathbb{F} \times \Omega \subseteq \mathbb{N} \times \mathbb{R}^n$.
- ▶ For each $m \in \mathbb{F}$ create Ω_m , a copy of Ω .
- ▶ Let Γ_m^\pm be respective exit and entrance boundaries into Ω_m (which can be artificially introduced if necessary), and suppose Q is a Markov kernel from $\Gamma^+ = \cup_{m \in \mathcal{F}} \Gamma_m^+$ to $\Gamma^- := \cup_{m \in \mathcal{F}} \Gamma_m^-$.
- ▶ While in some Ω_m^o , let the process undergo zig-zag dynamics with target $\pi(m, \cdot)$.
- ▶ When an exit boundary is hit, the state jumps with law Q and re-enters a different copy.

The zig-zag process on a hybrid state space

Take $\mathbb{F} = \{A, B\}$ and $\Omega = B(0, 1)$.



Piecewise deterministic Markov processes²

The generalised zig-zag process (m_t, x_t, v_t) is a PDMP with generator

$$Lf(m, x, v) := v \cdot \nabla_x f(m, x, v) + \sum_{i=1}^d \lambda_i(m, x, v) [f(m, x, F_i v) - f(m, x, v)]$$

whose domain $D(L)$ consists of measurable functions f satisfying

$$f(m, x, v) = \int_{(j, y, w) \in \Gamma^-} f(j, y, w) Q(m, x, v; j, dy, w)$$

for any $(m, x, v) \in \Gamma^+$, and some boundedness & continuity conditions.

²M Davis. Markov models and optimization, Chapman & Hall, 1993. Especially chapters 2 and 3.

Theorem³

Suppose the initial condition has a density, that \mathbb{F} is finite, that $\pi(m, \cdot) \in C^1(\Omega_m^o)$ for each $m \in \mathbb{F}$, that $\pi > 0$ on $\cup_{m \in \mathbb{F}} \Omega_m^o$, and

1. $\mathbb{E}_{(m,x,v)}[\#\text{jumps by time } t] < \infty$ for any (m, x, v) and $t > 0$,
2. for each $(x, v) \in \Gamma_m^+$ and $(y, w) \in \Gamma_j^-$,

$$\pi(m, x)Q(m, x, v; j, y, w) = \pi(j, y)Q(j, y, -w; m, x, -v),$$

3. for each $(x, v) \in \Gamma_m^+$,

$$\int_{(j,y,w) \in \Gamma^-} (w \cdot n(j, y))Q(m, x, v; j, dy, w) = -v \cdot n(m, x),$$

where $n(m, x)$ is the unit outward normal at $x \in \partial\Omega_m$,

4. and that Q is such that jumps facing corners are a null event.

Then π is a stationary distribution of (m_t, x_t, v_t) .

³J Koskela. Zig-zag sampling for discrete structures and non-reversible phylogenetic MCMC, arXiv:2004.08807.

Sketch proof

- ▶ The various technical assumptions guarantee that $C_b^1 \cap D(L)$ separates measures on the domain of the zig-zag process.
- ▶ The aim is to show $\mathbb{E}_\pi[Lf(m, x, v)] = 0$ for each $f \in C_b^1 \cap D(L)$.

$$\begin{aligned} & \mathbb{E}_\pi \left[v \cdot \nabla_x f(m, x, v) + \sum_{i=1}^d \lambda_i(m, x, v) [f(m, x, F_i v) - f(m, x, v)] \right] \\ &= \mathbb{E}_\pi [v \cdot \{ \nabla_x f(m, x, v) - f(m, x, v) \nabla_x \log \pi(m, x) \}] \\ &= \mathbb{E}_\pi [v \cdot \nabla_x f(m, x, v) - v \cdot \nabla_x f(m, x, v) - \text{Boundary terms}]. \end{aligned}$$

- ▶ The defining boundary property of the domain $D(L)$ and the two conditions on π and Q are exactly what you need to show that boundary terms vanish.

Ergodicity

If \mathbb{F} is finite and the process restricted to a single domain Ω_m with target $\pi(m, \cdot)$ is ergodic for each $m \in \mathbb{F}$, then the full process is ergodic provided

$$\int_{(x,v) \in \Gamma_m^+} \int_{(y,v) \in \Gamma_j^-} Q(m, x, v; j, dy, w) \pi(m, x) > 0$$

for enough ordered pairs $(m, j) \in \mathbb{F}^2$ to form a tour.

Some criteria for ergodicity of a single-domain process are known⁴.

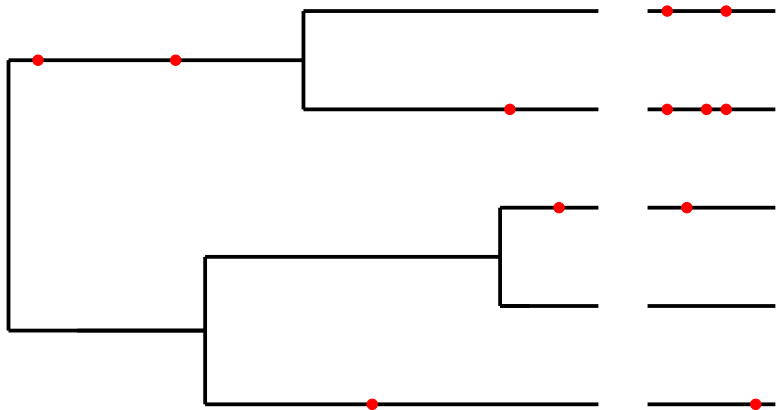
⁴J Bierkens, G Roberts and P-A Zitt. Ergodicity of the zig-zag process, Ann Appl Probab 29(4):2266–2301, 2019.

The coalescent⁵

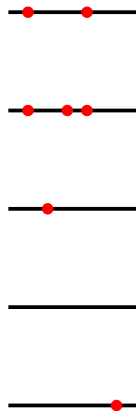
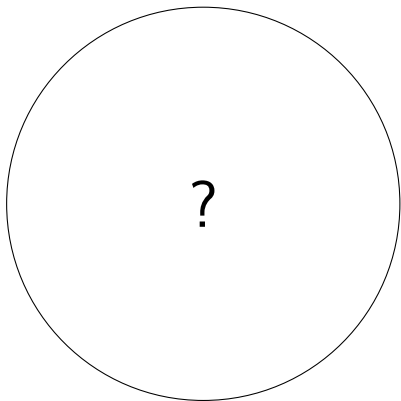
- ▶ $(\Pi_t)_{t \geq 0}$ with $\Pi_0 = \{\{1\}, \dots, \{n\}\}$.
- ▶ Holding times $T_k \sim \text{Exp}(\binom{k}{2})$.
- ▶ At jump times, two randomly chosen blocks merge.
- ▶ $i \sim j$ in $\Pi_t \Leftrightarrow i$ and j have a common ancestor at time t .
- ▶ Mutations as a Poisson point process with rate $\theta/2$ on the branches of the tree.

⁵J F C Kingman. The coalescent, Stoch Proc Appl 13(3):235–248, 1982.

The coalescent



The coalescent as missing data



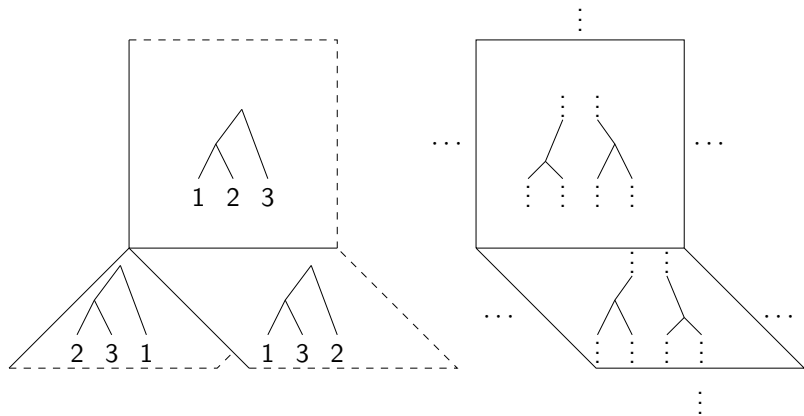
$$P(D|\theta) = \int_A P(D|A, \theta) P(A|\theta) dA$$

A geometric embedding of coalescent trees⁶

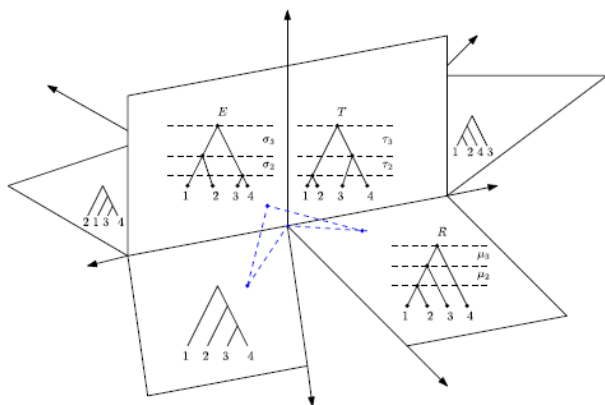
- ▶ Encode a tree as (E_n, \mathbf{t}_n) .
- ▶ $\mathbf{t}_n := (t_1, \dots, t_{n-1}) \in \mathbb{R}_+^{n-1}$ are the times between successive mergers.
- ▶ E_n is the *ranked topology*: an ordered set of nodes specifying the exact merger events.
- ▶ τ -space: separate copy of \mathbb{R}_+^{n-1} for each distinct E_n .
- ▶ Glue common faces of separate orthants together to form a connected space.
- ▶ $\#E_n = \frac{(n-1)!n!}{2^{n-1}}$.

⁶A Gavryushkin and A J Drummond. The space of ultrametric phylogenetic trees, J Theor Biol 403:197–208, 2016.

The three leaf τ -space



One third of the four leaf τ -space



⁶A Gavryushkin and A J Drummond. The space of ultrametric phylogenetic trees, J Theor Biol 403:197–208, 2016.

The infinite sites coalescent posterior

- ▶ Regard E_n as a set of edges.
- ▶ Given n observed sequences and a putative ancestral tree (E_n, \mathbf{t}_n) , for an edge $\gamma \in E_n$, define:
 - ▶ $m_\gamma :=$ number of mutations on γ .
 - ▶ $t_i \in \gamma$ if γ spans the i th interval.
 - ▶ $l_\gamma := \sum_{i:t_i \in \gamma} t_i$ be the branch length.

$$\pi(\theta, E_n, \mathbf{t}_n) \propto \left\{ \prod_{\gamma \in E_n} \frac{1}{m_\gamma!} \left(\frac{\theta l_\gamma}{2} \right)^{m_\gamma} \right\} \\ \times \exp \left(- \sum_{i=1}^{n-1} \frac{(n+1-i)(n+\theta-i)}{2} t_i \right).$$

Flip rates

$$\lambda_i(E_n, \mathbf{t}_n, \theta; \nu) := \nu_i \left(\frac{(n+1-i)(n+\theta-i)}{2} - \sum_{\gamma \in E_n: t_i \in \gamma} \frac{m_\gamma}{l_\gamma} \right) \vee 0,$$

$$\lambda_\theta(E_n, \mathbf{t}_n, \theta; \nu) := \nu_\theta \left(\sum_{i=1}^{n-1} \frac{n+1-i}{2} t_i - \frac{1}{\theta} \sum_{\gamma \in E_n} m_\gamma \right) \vee 0.$$

Important: $\nu_i = \pm 1 / \binom{n+1-i}{2}$.

Ranked topology crossings

- ▶ Exit and entrance boundaries := faces at which $t_i = 0$ or $\theta = 0$.
- ▶ Define the jump kernel Q via the orthants which intersect at a face.
- ▶ π is continuous at the boundaries of ranked topologies.
- ▶ Crossing into uniformly chosen adjacent topology when a boundary is hit satisfies the conditions on Q .
- ▶ So does reflecting at the $t_1 = 0$ and $\theta = 0$ boundaries.

Simulation study⁷

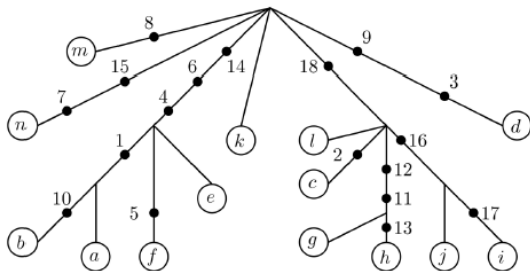


Figure 3.
Perfect phylogeny of the Griffiths and Tavaré (1994) data set.

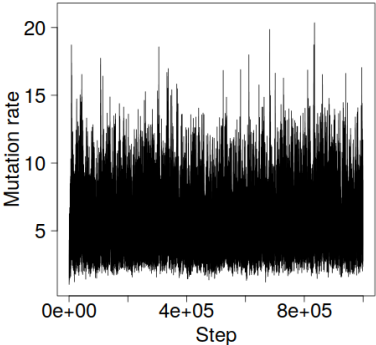
⁸A Hobolth, M K Uyenoyama and C Wiuf. Importance sampling for the infinite sites model, *Stat Appl Genet Mol Biol* 7(1) Article 32, 2008.

⁸R C Griffiths and S Tavaré. Ancestral inference in population genetics, *Stat Sci* 9:307–319, 1994.

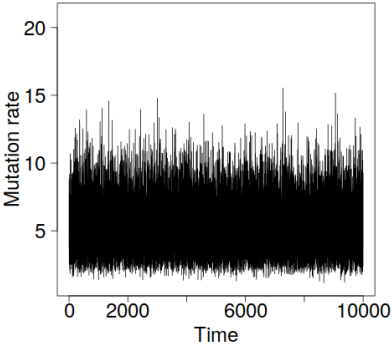
⁸R H Ward, B L Frazier, K Dew and S Pääbo. Extensive mitochondrial diversity within a single Amerindian tribe, *Proc Natl Acad Sci USA* 88:8720–8724, 1991.

Mutation rate

Metropolis

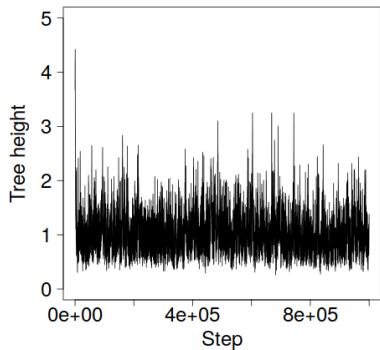


Zig-zag

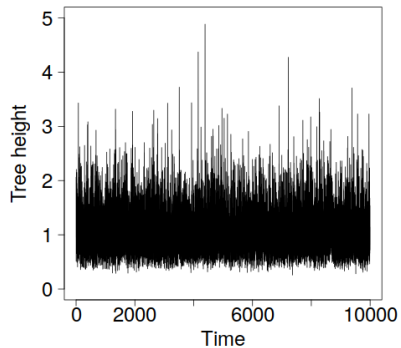


Tree height

Metropolis

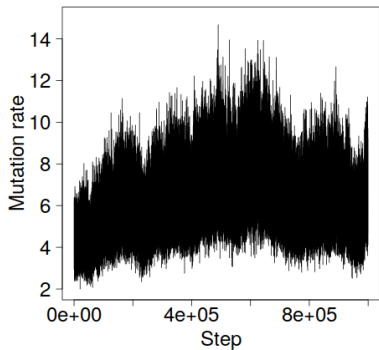


Zig-zag

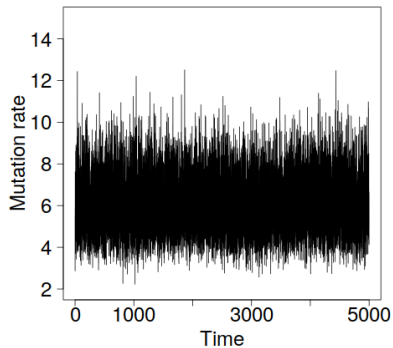


Mutation rate, sample size = 550

Metropolis

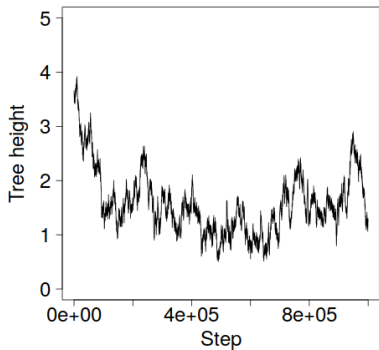


Zig-zag

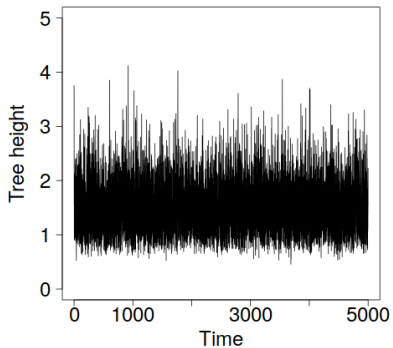


Tree height, sample size = 550

Metropolis

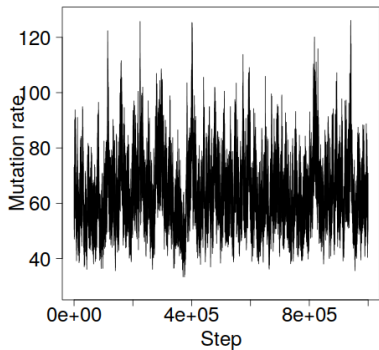


Zig-zag

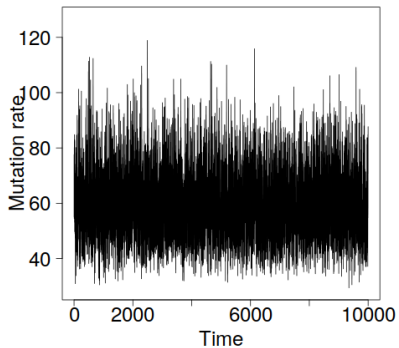


Mutation rate, $\theta = 55$

Metropolis

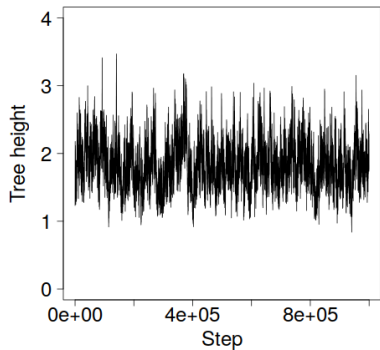


Zig-zag

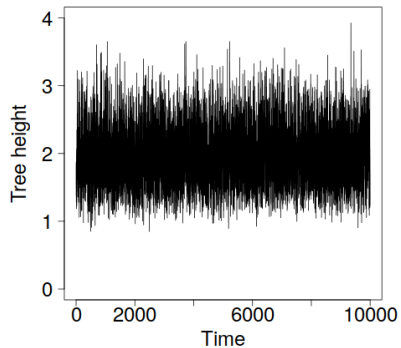


Tree height, $\theta = 55$

Metropolis



Zig-zag



What about a nuclear reactor?

